AMENDMENTS TO THE CLAIMS

1. (Currently Amended) A method of correlating ~~the~~ an FcαRI induced function of intracellular calcium flux or interleukin-6 release or TNFα release of a cell in a host expressing FcαRI ~~and cellular susceptibility to a disease~~ with an FcαRI amino acid sequence, said method comprising:

identifying [[a]] said FcαRI genotype at nucleotide 844 corresponding to an amino acid sequence codon 248 of said cell ~~for FcαRI alleles selected from the group consisting of: FcαRIa 87R/87R, FcαRIA 92D/92N, FcαRIa 132F/132L, FαRI 245P/245L and FcαRI 248S/248G~~ as being either glycine or serine;

quantifying [[an]] said FcαRI induced function ~~selected from the group consisting of: FCαRI specific phagocytosis, oxidative burst and cytokine production~~ by said cell expressing ~~said~~ FcαRI ~~genotype~~; and

comparing FcαRI induced function by said cell and FcαRI induced function by a second cell, said second cell expressing a second FcαRI genotype at nucleotide 844 that corresponds to an alternate to said amino acid sequence codon 248~~, wherein correlation of the FcαRI induced function and cellular susceptibility to disease is indicated by a difference in FcαRI induced function detected by said comparing~~, wherein the FcαRI induced function of intracellular calcium flux or interleukin-6 release or TNFα release of the cell in the host expressing FcαRI correlates with the FcαRI amino acid sequence codon 248.

Claims 2-11 (Canceled)

12. (Original) The method of claim 1 wherein said cell is selected from the group consisting of: a neutrophil, a monocyte, a myeloid cell, and a mucus secreting cell.

13. (Currently Amended) A method for determining FcαRI ~~alleles~~ induced function of increased intracellular calcium flux or greater interleukin-6 release or less TNFα release specific to an individual human, said method comprising: genotyping DNA encoding FcαRI for ~~a polymorphism affecting an FcαRI induced function selected from the group consisting of: FcαRIa 87R/87R, FcαRIA 92D/92N, FcαRIa 132F/132L, FαRI 245P/245L and FcαRI 248S/248G~~ nucleotide 844 corresponding to a codon identity at codon 248 as being glycine, said DNA being obtained from said individual human, wherein the FcαRI induced function specific to said individual human is relative to codon 248 being serine.

Claims 14-19 (Canceled)

20. (Original) The method of claim 13 wherein genotyping utilizes PCR typing with a sequence specific primer for a polymorphic exon.

21. (Previously Presented) The method of claim 20 wherein said primer is selected from the group consisting of SEQ ID No. 1, SEQ ID No. 2, SEQ ID No. 3, or SEQ ID No. 4.

22. (Withdrawn) A method for correlating the ability of a cell to bind IgA, and cellular susceptibility to a disease, said method comprising:

identifying a FcαRI phenotype of said cell;

quantifying IgA binding by said cell; and

comparing IgA binding by said cell to that of a second cell, said second cell having a second phenotype FcαRI.

23. (Withdrawn) The method of claim 22 wherein identifying said FcαRI phenotype utilizes amino acid sequencing.

24. (Withdrawn) The method of claim 22 wherein identifying said FcαRI phenotype utilizes glycosylate characterization.

25. (Withdrawn) The method of claim 22 wherein identifying said FcαRI phenotype utilizes antibody binding.

26. (Currently Amended) A method of prognosticating a human ~~immunoresponse to a disease~~ CD89 expressing cellular response, said method comprising:

establishing a correlation between a FcαRI genotype ~~for a FcαRI alleles selected from the group consisting of: FcαRIa 87R/87R, FcαRIA 92D/92N, FcαRIa 132F/132L, FαRI 245P/245L and FcαRI 248S/248G and clinical outcome of said disease~~ at nucleotide 844 as to A and a cellular response selected from the group consisting of: increased intracellular calcium flux, greater interleukin-6 release and less TNFα release relative to nucleotide 844 being G;

genotyping a patient for FcαRI to yield a patient FcαRI genotype at nucleotide 844;

comparing said FcαRI genotype with said patient genotype; and

determining ~~clinical outcome for said patient~~ said cellular response based on said patient genotype~~, wherein determining said clinical outcome is indicative of a human immunoresponse to a disease~~, wherein the human CD89 expressing cellular response is prognosticated based on the nucleotide 844 genotype being A and the cellular response is selected from the group consisting of: increased intracellular calcium flux, greater interleukin-6 release and less TNFα release.

27. (Original) The method of claim 26 wherein genotyping utilizes PCR typing with a sequence specific primer for a polymorphic exon.

28. (Original) The method of claim 27 wherein said primer is selected from the group consisting of those shown in SEQ ID Nos. 1, 2, 3 and 4.

29. (Original) The method of claim 26 wherein genotyping comprises purifying FcαRI expressing cells from said patient; extracting nucleic acids from said cells; and determining whether the nucleic acid encodes a predetermined polymorphic FcαRI nucleic acid sequence.

30. (Original) The method of claim 29 wherein the nucleic acid is selected from the group consisting of: RNA and DNA.

Claims 31-46 (Canceled)